

**Amendments to the Claims:**

This listing of claims will replace all prior versions, and listings, of claims in the application:

**Listing of Claims:**

1. (currently amended) A metabolically engineered micro-organism which is yeast having an operative first metabolic pathway in which a first metabolite is transformed into a second metabolite in a reaction in which NAD is a cofactor for a first enzyme, said reaction step producing NADH, and in which said second metabolite is transformed into at least one further metabolite in a reaction catalysed by a second enzyme, and having an operative second metabolic pathway characterized by an enzyme activity in excess of a native level in respect of a third enzyme catalyzing a non-reversible reaction in which NADP is a cofactor and NADPH is a product and in which said first metabolite is transformed into a said further metabolite without the involvement of said second enzyme.

2. (original) A micro-organism as claimed in claim 1, wherein said first metabolic pathway is a native pathway.

3. (currently amended) A micro-organism as claimed

| ~~in any preceding claim 1~~, wherein said first enzyme is a  
phosphorylating dehydrogenase.

4. (currently amended) A micro-organism as claimed

| in claim 1~~or claim 2~~, wherein said second enzyme is a kinase.

5. (original) A micro-organism as claimed in claim  
3, wherein said third enzyme is a non-phosphorylating  
dehydrogenase.

6. (original) A micro-organism as claimed in claim  
5, wherein said third enzyme is GAPN (EC 1.2.1.9).

7. (original) A micro-organism as claimed in claim  
6, wherein said first enzyme is GAPDH (EC 1.2.1.12).

8. (currently amended) A micro-organism as claimed  
| ~~in any preceding claim 1~~, wherein at least one copy of a  
genetic sequence encoding said third enzyme has been  
recombinantly introduced into said organism.

9. (currently amended) A micro-organism as claimed  
in ~~any preceding claim 1~~, wherein a genetic sequence encoding  
said third enzyme is operatively linked to an expression  
signal not natively associated with said genetic sequence in  
said organism.

10. (deleted)

11. (currently amended) A micro-organism as claimed  
in claim 10, which is a ~~micro-organism yeast~~ belonging to the  
genus *Saccharomyces*, *Klyuveromyces*, *Candida*, *Pichia*,  
*Debaromyces*, *Hansenula*, *Yarrowia*, *Zygosaccharomyces* or  
*Schizosaccharomyces*.

12. (currently amended) A micro-organism as claimed  
in claim ~~10~~11, which is a strain of *Saccharomyces cerevisiae*,  
*S. kluyveri*, *S. bayanus*, *S. exiguum*, *S. sevazzi*, *S. uvarum*,  
*Klyuveromyces lactis* *K. marxianus* var. *marxianus*, *K.*  
*thermotolerans*, *Candida utilis* *C. tropicalis*, *Pichia stipidis*,  
*P. pastoris*, *P. sorbitophila*, *Debaromyces hansenii*, *Hansenula*  
*polymorpha*, *Yarrowia lipolytica*, *Zygosaccharomyces rouxii* or  
*Schizosaccharomyces pombe*.

13. (currently amended) A genetically transformed micro-organism which is a yeast containing one or more copies of a heterologous DNA sequence encoding GAPN (EC 1.2.1.9) operatively associated with an expression signal and having a functional native or heterologous expression capability for GAPDH (EC 1.2.12).

14. (currently amended) A method of producing a desired metabolic product with decreased production of an undesired metabolic product, comprising culturing a micro-organism as claimed in ~~any preceding claim 1 wherein the~~ desired metabolic product is the product of the reaction catalyzed by said third enzyme, and the undesired metabolic product is the product of the reaction catalyzed by the second enzyme.

15. (original) A method as claimed in claim 14, wherein the desired product is ethanol, lactic acid, citric acid, an amino acid or an antibiotic.

16. (currently amended) A method as claimed in claim 14 or ~~claim 15~~, wherein said undesired metabolic product is glycerol, acetate or an amino acid.